pD17-cJ-dCH2.H1

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        10          20          30          40          50          60          70          80          90
GACGGATCGG  GAGATCTGCT  AGGTGACCTG  AGGCGCGCCG  GCTTCGAATA  GCCAGAGTAA  CCTTTTTTTT  TAATTTTATT  TTATTTTATT
CTGCCTAGCC  CTCTAGACGA  TCCACTGGAC  TCCGCGCGGC  CGAAGCTTAT  CGGTCTCATT  GGAAAAAAAA  ATTAAAATAA  AATAAAATAA

       100         110         120         130         140         150         160         170         180
TTTGAGATGG  AGTTTGGCGC  CGATCTCCCG  ATCCCCTATG  GTCGACTCTC  AGTACAAATCT  GCTCTGATGC  CGCATAGTTA  AGCCAGTATC
AAACTCTACC  TCAAACCGCG  GCTAGAGGGC  TAGGGGATAC  CAGCTGAGAG  TCATGTTTAGA  CGAGACTACG  GCGTATCAAT  TCGGTCATAG

       190         200         210         220         230         240         250         260         270
TGCTCCCTGC  TTGTGTGTTG  GAGGTGCTG   AGTAGTGCGC  GAGCAAAATT  TAAGCTACAA  CAAGGCAAGG  CTTGACCGAC  AATTGCATGA
ACGAGGGACG  AACACACAAC  CTCCAGCGAC  TCATCACGCG  CTCGTTTTAA  ATTCGATGTT  GTTCCGTTCC  GAACTGGCTG  TTAACGTACT

       280         290         300         310         320         330         340         350         360
AGAATCTGCT  TAGGGTTAGG  TGCTTCGCGA  CGTTTGCGC   TGTACGGGCC  AGATATACGC  GTTGACATTG  ATTTATTGACT  AGTTATTAAT
TCTTAGACGA  ATCCCAATCC  ACGAAGCGCT  GCAAAACGCG  ACATGCCCGG  TCTATATGCG  CAACTGTAAC  TAATAACTGA  TCAATAATTA

       370         380         390         400         410         420         430         440         450
TACGGGGTCA  TTAGTTCATA  GCCCATATAT  GGAGTTCCGC  GTTACATAAC  GTTACGGTAAA  TTACGGTAAA  TGGCCCGCCT  GGCTGACCGC
ATGCCCCAGT  AATCAAGTAT  CGGGTATATA  CCTCAAGGCG  CAATGTATTG  AATGCCATTT  AATGCCATTT  ACCGGGCGGA  CCGACTGGCG

       460         470         480         490         500         510         520         530         540
CCAACGACCC  ACGTCAATAA  TGACGTATGT  TCCCATAGTA  ACGCCAATAG  GGACTTTCCA  TTGACGTCAA  TGGGTGGACT
GGTTGCTGGG  TGCAGTTATT  ACTGCATACA  AGGGTATCAT  TGCGGTTATC  CCTGAAAGGT  AACTGCAGTT  ACCCACCTGA

       550         560         570         580         590         600         610         620         630
ATTACGGTA   TTGGCAGTAC  ATCAAGTGTA  TCATATGCCA  AGTACGCCCC  CTATTGACGT  CAATGACGGT  AAATGGCCCG
TAAATGCCAT  AACCGTCATG  TAGTTCACAT  AGTATACGGT  TCATGCGGGG  GATAACTGCA  GTTACTGCCA  TTTACCGGGC

       640         650         660         670         680         690         700         710         720
CCTGGCATTA  ATGACCTTAT  GGGACTTTCC  TACTTGGCAG  TACATCTACG  TATTAGTCAT  CGCTATTTACC  ATGGTGATGC
GGACCGTAAT  ACGGGTCATG  CCCTGAAAGG  ATGAACCGTC  ATGTAGATGC  ATAATCAGTA  GCGATAATGG  TACCACTACG

       730         740         750         760         770         780         790         800         810
GGTTTGGCA   GTACATCAAT  GGGCGTGGAT  AGCGGTTTGA  TTTCCAAGTC  TCCACCCCAT  TGACGTCAAT  GGGAGTTTGT
CCAAAACCGT  CATGTAGTTA  CCCGCACCTA  TCGCCAAACT  AAAGGTTCAG  AGGTGGGGTA  ACTGCAGTTA  CCCTCAAACA

       820         830         840         850         860         870         880         890         900
TTTGGCACCA  GACTTTCCAA  AATGTCGTAA  CAACTCCGCC  CCATTGACGC  TAGGCGTGTA  CGGTGGGAGG
AAACCGTGGT  CTGAAAGGTT  TTACAGCATT  GTTGAGGCGG  GGTAACTGCG  ATCCGCACAT  GCCACCCCTCC
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Figure 14A
(SEQ ID NO.: 10 – Primary Sequence)
(SEQ ID NO.: 28 – Complement)

pD17-cJ-dCH2.H1

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910                 920                 930                 940                 950                 960                 970                 980                 990
TCTATATAAG          CAGAGCTCTC          TGGCTAACTA          GAGAACCCAC          TGCTTACTGG          CTTATCGAAA          TTAATACGAC          TCACTATAGG          GAGACCCAAG
AGATATATTC          GTCTCGAGAG          ACCGATTGAT          CTCTTGGGTG          ACGAATGACC          GAATAGCTTT          AATTATGCTG          AGTGATATCC          CTCTGGGTTC

1000                1010                1020                1030                1040                1050                1060                1070                1080
CTTGGTACCA          ATTTAAATTG          ATATCTCCTT          AGGTCTCGAG          TCTCTAGATA          ACCGGTCAAT          CGATTGGAAT          TCTTGCGGCC          GCTTGCTAGC
GAACCATGGT          TAAATTTAAC          TATAGAGGAA          TCCAGAGCTC          AGAGATCTAT          TGGCCAGTTA          GCTAACC-TA          AGAACGCCGG          CGAACGATCG

1090                1100                1110                1120                1130                1140                1150                1160                1170
CACCATGGAG          TTGTGGTTAA          GCTTGGTCCT          TCCTTGTCCT          TGTTTTAAAA          GGTGTCCAGT          GTGAAGTGAA          TCTGGTGGAG          TCTGGGGGAG
GTGGTACCTC          AACACCAATT          CGAACCAGGA          AGGAACAGGA          ACAAAATTTT          CCACAGGTCA          CACTTCACTT          AGACCACCTC          AGACCCCCTC

1180                1190                1200                1210                1220                1230                1240                1250                1260
GCTTAGTGCA          GCCTGGAGGG          TCCCTGAAAG          TCTCCTGTGT          AACCTCTCGA          TTCACTTTCA          GTGACTATTA          CATGTATTGG          GTTCGCCAGA
CGAATCACGT          CGGACCTCCC          AGGGACTTTC          AGAGGACACA          TTGGAGACCT          AAGTGAAAGT          CACTGATAAT          GTACATAACC          CAAGCGGTCT

1270                1280                1290                1300                1310                1320                1330                1340                1350
CTCCAGAGAA          GAGGCTGGAG          TGGGTCGCAT          ACATTAGTCA          AGGTGGTGAT          ATAACCGACT          ATCCAGACAC          TGTAAAGGGT          CGATTCACCA
GAGGTCTCTT          CTCCGACCTC          ACCCAGCGTA          TGTAATCAGT          TCCACCACTA          TATTGGCTGA          TAGGTCGTGTG         ACATTTCCCA          GCTAAGTGGT

1360                1370                1380                1390                1400                1410                1420                1430                1440
TCTCCAGAGA          CAATGCCAAG          AACCCCTGT           ACCTGCAAAT          GAGCCGTCTG          AAGTCTGAGG          ACACAGCCAT          GTATTACTGT          GCAAGAGGCC
AGAGGTCTCT          GTTACGGTTC          TTGTGGGACA          TGGACGTTTA          CTCGGCAGAC          TTCAGACTCC          TGTGTCGGTA          CATAATGACA          CGTTCTCCGG

1450                1460                1470                1480                1490                1500                1510                1520                1530
GGCCTGGTTT          GCTTACTGGG          GCCAAGGGAC          TCTGGTCACG          GTCTCTGTAG          CTAGCACCAA          GGGCCCATCG          GTCTTCCCCC          GTGGTCACCG
CCGGACCAAA          CGAATGACCC          CGGTTCCCTG          AGACCAGTGC          CAGAGACATC          GATCGTGGTT          CCCGGGTAGC          CAGAGAGGGG          CACCAGTGGC

1540                1550                1560                1570                1580                1590                1600                1610                1620
TGGCACCCTC          CTCCAAGAGC          ACCTCTGGGG          GCACAGCGGC          CCTGGGCTGC          CTGGTCAAGG          ACTACTTCCC          CGAACCGGTG          ACGGTGTCGT
ACCGTGGGAG          GAGGTTCTCG          TGGAGACCCC          CGTGTCGCCG          GGACCCGACG          GACCAGTTCC          TGATGAAGGG          GCTTGGCCAC          TGCCACAGCA

1530                1640                1650                1660                1670                1680                1690                1700                1710
GGAACTCAGG          CGCCCTGACC          AGCGGCGTGC          ACACCTTCCC          GGCTGTCCTA          CAGTCCTCAG          GACTCCTACTC         CCTCAGCAGC          GTGGTCACCG
CCTTGAGTCC          GCGGGACTGG          TCGCCGCACG          TGTGGAAGGG          CCGACAGGAT          GTCAGGAGTC          CTGAGATGAG          GGAGTCGTCG          CACCAGTGGC

1720                1730                1740                1750                1760                1770                1780                1790                1800
TGCCCTCCCAG         CAGCTTGGGC          ACCCAGACCT          ACATCTGCAA          CGTGAATCAC          AAGCCCAGCA          ACACCAAGGT          GGACAAGAAA          GTTGGTGAGA
ACGGGAGGTC          GTCGAACCCG          TGGGTCTGGA          TGTAGACGTT          GCACTTAGTG          TTCGGGTCGT          TGTGGTTCCA          CCTGTTCTTT          CAACCACTCT
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Figure 14B
(SEQ ID NO.: 10 – Primary Sequence)
(SEQ ID NO.: 28 – Complement)

pD17-cJ-dCH2.H1

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1810 GGCCAGCACA GGGAGGGAGG GTGTCTGCTG GAAGCCAGGC TCAGGCGTCC TGCCTGGACG CATCCCGGCT ATGCAGCCCC AGTCCAGGGC
     CCGGTCGTGT CCCTCCCTCC CACAGACGAC CTTCGGTCCG AGTCCGCAGG ACGGACCTGC GTAGGGCCGA TACGTCGGGG TCAGGTCCCG

1900 AGCAAGGCAG GCCCCGTCTG CCTCTTCACC CGGAGGCCTC TGCCCGCCCC ACTCATGCTC AGGGAGAGGG TCTTCTGGCT TTTTCCCCAG
     TCGTTCCGTC CGGGGCAGAC GGAGAAGTGG GCCTCCGGAG ACGGGCGGGG TGAGTACGAG TCCCTCTCCC AGAAGACCGA AAAAGGGGTC

1990 GCTCTGGGCA GGCACAGGCT AACCCAGGCC CTGCACACAA AGGGGCAGGT GCTGGGCTCA GACCTGCCAA GAGCCATATC
     CGAGACCCGT CCGTGTCCGA TTGGGTCCGG GACGTGTGTT TCCCCGTCCA CGACCCGAGT CTGGACGGTT CTCCGTATAG

2080 CGGGAGGACC CTGCCCCTGA CCTAAGCCCA CCCCAAAGGC CAAACTCTCC ACTCCCTCAG CTCGGACACC TTCTCTCCTC CCAGATTCCA
     GCCCTCCTGG GACGGGGACT GGATTCGGGT GGGGTTTCCG GTTTGAGAGG TGAGGGAGTC GAGCCTGTGG AAGAGAGGAG GGTCTAAGGT

2170 ATCTTCTCTC TGCAGAGCCC AAATCTTGTG ACAAAACTCA CACATGCCCA CCGTGCCCAG GTAAGCCAGC CCAGGCCTCG GGTCCGGAGC
     TAGAAGAGAG ACGTCTCGGG TTTAGAACAC TGTTTTGAGT GTGTACGGGT GGCACGGGTC CATTCGGTCG GGTCCGGAGC CCAGGCCTCG

2260 CAAGGCGGGA CCCTCCAGCT CAGGTGCCCT AGAGTAGCCT GCATCCAGGG ACACACCACG TGGGTACCAA CATGTCCGGA GCCACATGGA
     GTTCCGCCCT GGGAGGTCGA GTCCACGGGA TCTCATCGGA CGTAGGTCCC TGTGTGGTGC ACCCATGGTT GTACAGGCCT CGGTGTACCT

2350 CTCGGCCCAC CCTCTGCCCT GAGAGTGACC GCTGTACCAA CCTCTGTCCC TACAGGGCAG CCCGAGAAAC CACAGGTGTA
     GAGCCGGGTG GGAGACGGGA CTCTCACTGG CGACATGGTT GGAGACAGGG ATGTCCCGTC GGGCTCTTTG GTGTCCACAT

2440 CACCCTGCCC CCATCCCGGG ATGAGCTGAC CAAGAACCAG GTCAGCCTGA CCTGCCTGGT CAAAGGCTTC TATCCCAGCG ACATCGCCGT
     GTGGGACGGG GGTAGGGCCC TACTCGACTG GTTCTTGGTC CAGTCGGACT GGACGGACCA GTTTCCGAAG ATAGGGTCGC TGTAGCGGCA

2530 GGAGTGGGAG AGCCGGAGAA CAACTACAAG ACCACGCCTC CCGTGCTGGA CTCCGACGGC TCCTTCTTCC TCTACAGCAA
     CCTCACCCTC TCGGCCTCTT GTTGATGTTC TGGTGCGGAG GGCACGACCT GAGGCTGCCG AGGAAGAAGG AGATGTCGTT

2620 GCTCACCGTG GACAAGAGCA GGTGGCAGCA GGGGAACGTC TTCTCATGCT CCGTGATGCA TGAGGCTCTG CACAACCACT ACACGCAGAA
     CGAGTGGCAC CTGTTCTCGT CCACCGTCGT CCCCTTGCAG AAGAGTACGA GGCACTACGT ACTCCGAGAC GTGTTGGTGA TGTGCGTCTT
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Figure 14C
(SEQ ID NO.: 10 – Primary Sequence)
(SEQ ID NO.: 28 – Complement)

pD17-cJ-dCH2.H1

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2710 GAGCCTCTCC   2720 CTGTCTCCGG   2730 GTAAATGAGT   2740 GCGACGGCCG   2750 GCAAGCCCCC
     CTCGGAGAGG        GACAGAGGCC        CATTTACTCA        CGCTGCCGGC        CGTTCGGGGG
2760 GCTCCCGGGG   2770 CTCTCGCGGT   2780 CGCACGAGGA   2790 TGCTTGGCAC
     CGAGGGCCCC        GAGAGCGCCA        GCGTGCTCCT        ACGAACCGTG

2800 GTACCCCCTG   2810 TACATACTTC   2820 CCGGGGCGCC   2830 AGGATGGAAA   2840 TAAAGCACCC
     CATGGGGGAC        ATGTATGAAG        GGCCCGCGGG        TCGTACCTTT        ATTTCGTGGG
2850 TGGCCCCCTG   2860 AGCGGCTGCC   2870 CGAGACTGTG   2880 ATGGTTCTTT
     ACCCGGGGAC        TCGCGACGGG        GCTCTGACAC        TACCAAGAAA

2890 CCACGGGTCA   2900 GGCCGAGTCT   2910 GAGGCCTGAG   2920 TGGCATGAGG   2930 GAGGCAGAGC
     GGTGCCCAGT        CCGGCTCAGA        CTCCGGACTC        ACCGTACTCC        CTCCGTCTCG
2940 GGGTCCCACT   2950 GTCCCCACAC   2960 TGGCCCAGGC   2970 TGTGCAGGTG
     CCCAGGGTGA        CAGGGGTGTG        ACCGGGTCCG        ACACGTCCAC

2980 TGCCTGGGCC   2990 CCCTAGGGTG   3000 GGGCTCAGCC   3010 AGGGGCTGCC   3020 CTCGGCAGGG
     ACGGACCCGG        GGGATCCCAC        CCCGAGTCGG        TCCCCGACGG        GAGCCGTCCC
3030 TGGGGGATTT   3040 GCCAGCGTGG   3050 CCCTCCCTCC   3060 AGCAGCACCT
     ACCCCCTAAA        CGGTCGCACC        GGGAGGGAGG        TCGTCGTGGA

3070 GCCCTGGGCT   3080 GGGCCACGGG   3090 AAGCCCTAGG   3100 AGCCCCTGGG   3110 GACAGACACA
     CGGGACCCGA        CCCGGTGCCC        TTCGGGATCC        TCGGGGACCC        CTGTCTGTGT
3120 CAGCCCCTGC   3130 CTCTGTAGGA   3140 GACTGTCCTG   3150 TTCTGTGAGC
     GTCGGGGACG        GAGACATCCT        CTGACAGGAC        AAGACACTCG

3160 GCCCCTGTCC   3170 TCCCGACCTC   3180 CATGCCCACT   3190 CGGGGCCATG   3200 CCTAGTCCAT
     CGGGGACAGG        AGGGCTGGAG        GTACGGGTGA        GCCCCGGTAC        GGATCAGGTA
3210 GTGCGTAGGG   3220 ACAGGCCCTC   3230 CCTCACCCAC   3240 CTACCCCCAC
     CACGCATCCC        TGTCCGGGAG        GGAGTGGGTG        GATGGGGGTG

3250 CCGTGATTGG   3260 CCTGGCTGCC   3270 CTGCCCAGCC   3280 TCGCACCGGC   3290 ATGGGGACAC
     GGCACTAACC        GGACCGACGG        GACGGGTCGG        AGCGTGGCCG        TACCCCTGTG
3300 AACCGACTCC   3310 GGGGACATGC   3320 ACTCTCGGGC   3330 CCTGTGGAGG
     TTGGCTGAGG        CCCCTGTACG        TGAGAGCCCG        GGACACCTCC

3340 GACTGGTGCA   3350 GATGCCCACA   3360 CACACACTCA   3370 GCCCAGACCC   3380 GTTCAACAAA
     CTGACCACGT        CTACGGGTGT        GTGTGTGAGT        CGGGTCTGGG        CAAGTTGTTT
3390 CCCCGCACTG   3400 AGGTTGGCCG   3410 GCCACACGGC   3420 CACCACACAC
     GGGGCGTGAC        TCCAACCGGC        CGGTGTGCCG        GTGGTGTGTG

3430 ACACGTGCAC   3440 GCCTCACACA   3450 CGGAGCCTCA   3460 CCCGGGCGAA   3470 CTGCACAGCA
     TGTGCACGTG        CGGAGTGTGT        GCCTCGGAGT        GGGCCCGCTT        GACGTGTCGT
3480 CCCAGACCAG   3490 AGCAAGGTCC   3500 TCGCACACGT   3510 GAACACTCCT
     GGGTCTGGTC        TCGTTCCAGG        AGCGTGTGCA        CTTGTGAGGA

3520 CGGACACAGG   3530 CCCCCACGAG   3540 CCCCACGCGG   3550 CACCTCAAGG   3560 CCCACGAGCC
     GCCTGTGTCC        GGGGGTGCTC        GGGGTGCGCC        GTGGAGTTCC        GGGTGCTCGG
3570 TCTCGGCAGC   3580 TTCTCCACAT   3590 GCTGACCTGC   3600 TCAGACAAAC
     AGAGCCGTCG        AAGAGGTGTA        CGACTGGACG        AGTCTGTTTG
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Figure 14D
(SEQ ID NO.: 10 – Primary Sequence)
(SEQ ID NO.: 28 – Complement)

pD17-cJ-dCH2.H1

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   3610          3620          3630          3640          3650          3660          3670          3680          3690
CCAGCCCTCC    TCTCACAAGG    GTGCCCCTGC    AGCCGCCACA    CACACACAGG    GGATCACACA    CCACGTCACG    TCCCTGGCCC    TGGCCCACTT
GGTCGGGAGG    AGAGTGTTCC    CACGGGGACG    TCGGCGGTGT    GTGTGTGTCC    CCTAGTGTGT    GGTGCAGTGC    AGGGACCGGG    ACCGGGTGAA

   3700          3710          3720          3730          3740          3750          3760          3770          3780
CCCAGTGCCG    CCCTTCCCTG    CAGGACGGAT    CAGCCTCGAC    TGTGCCTTCT    AGTTGCCAGC    CATCTGTTGT    TTGCCCCTCC    CCCGTGCCTT
GGGTCACGGC    GGGAAGGGAC    GTCCTGCCTA    GTCGGAGCTG    ACACGGAAGA    TCAACGGTCG    GTAGACAACA    AACGGGGAGG    GGGCACGGAA

   3790          3800          3810          3820          3830          3840          3850          3860          3870
CCTTGACCCT    GGAAGGTGCC    ACTCCCACTG    TCCTTCCCTA    ATAAAATGAG    GAAATTGCAT    CGCATTGTCT    GAGTAGGTGT    CATTCTATTC
GGAACTGGGA    CCTTCCACGG    TGAGGGTGAC    AGGAAAGGAT    TATTTTACTC    CTTTAACGTA    GCGTAACAGA    CTCATCCACA    GTAAGATAAG

   3880          3890          3900          3910          3920          3930          3940          3950          3960
TGGGGGTGG     GGTGGGGCAG    GACAGCAAGG    GGGAGGATTG    GGAAGACAAT    AGCAGGCATG    CTGGGGATGC    GGTGGGCTCT    ATGGCTTCTG
ACCCCCACC     CCACCCCGTC    CTGTCGTTCC    CCCTCCTAAC    CCTTCTGTTA    TCGTCCGTAC    GACCCCTACG    CCACCCGAGA    TACCGAAGAC

   3970          3980          3990          4000          4010          4020          4030          4040          4050
AGGCGGAAAG    AACCAGCTGG    GGCTCTAGGG    GGTATCCCCA    CGGCCCCTGT    AGCGGGCGCAT   TAAGCGCGGC    GGTGTGGTG     GTTACGGCA
TCCGCCTTTC    TTGGTCGACC    CCGAGATCCC    CCATAGGGGT    GCCGGGGACA    TCGCCCGCGTA   ATTCGCGCCG    CCCACACCAC    CAATGCCGGT

   4060          4070          4080          4090          4100          4110          4120          4130          4140
GGCTGACCGC    TACACTTGCC    AGCGCCCTAG    CGCCCGCTCC    TTTCGCTTTC    TTCCCTTCCT    GTTCGCCGGG    CCTTCTCAAAA   CCTCTCAAAA
CGCACTGGCG    ATGTGAACGG    TCGCGGGATC    GCGGGCGAGG    AAAGCGAAAG    AAGGGAAGGA    CAAGCGGCCC    GGAGAGTTTT    GGAGAGTTTT

   4150          4160          4170          4180          4190          4200          4210          4220          4230
AAGGGAAAAA    AAGCATGCAT    CTCAATTAGT    CAGCAACCAT    AGTCCCGCCC    CTAACTCCGC    CCATCCCGCC    CCTAACTCCG    CCCAGTTCCG
TTCCCTTTTT    TTCGTACGTA    GAGTTAATCA    GTCGTTGGTA    TCAGGGCGGG    GATTGAGGCG    GGTAGGGCGG    GGATTGAGGC    GGGTCAAGGC

   4240          4250          4260          4270          4280          4290          4300          4310          4320
CCCATTCTCC    GCCCCATGGC    TGACTAATTT    TTTTTATTTA    TGGCAGAGGCC   GAGGCCGCCT    CGGCCTCTGA    GCTATTCCAG    AAGTAGTGAG
GGGTAAGAGG    CGGGGTACCG    ACTGATTAAA    AAAAATAAAT    ACGTCTCCGG    CTCCGGCGGA    GCCGGAGACT    CGATAAGGTC    TTCATCACTC

   4330          4340          4350          4360          4370          4380          4390          4400          4410
GAGGCTTTTT    TGGAGGCCTA    GGCTTTTGCA    AAAAGCTTGG    ACAGCTCAGG    GCTGCGATTT    CGCGCCAAAC    TTGACGGCAA    TCCTAGCGTG
CTCCGAAAAA    ACCTCCGGAT    CCGAAAACGT    TTTTCGAACC    TGTCGAGTCC    CGACGCTAAA    GCGCGGTTTG    AACTGCCGTT    AGGATCGCAC

   4420          4430          4440          4450          4460          4470          4480          4490          4500
AAGGCTGGTA    GGATTTTATC    CCCGCTGCCA    TCATGGTTCG    ACCATTGAAC    TGCATCGTCG    CCGTGTCCCA    AAATATGGGG    ATTGGCAAGA
TTCCGACCAT    CCTAAAATAG    GGGCGACGGT    AGTACCAAGC    TGGTAACTTG    ACGTAGCAGC    GGCACAGGGT    TTTATACCCC    TAACCGTTCT
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Figure 14E
(SEQ ID NO.: 10 – Primary Sequence)
(SEQ ID NO.: 28 – Complement)

pD17-cJ-dCH2.H1

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4510 ACGGAGACCT   4520 ACCCTGGCCT   4530 CCGCTCAGGA   4540 ACGAGTTCAA   4550 GTACTTCCAA
     TGCCTCTGGA        TGGGACCGGA        GGCGAGTCCT        TGCTCAAGTT        CATGAAGGTT
4560 AGAATGACCA   4570 CAACCTCTTC   4580 AGTGGAAGGT   4590 AAACAGAATC
     TCTTACTGGT        GTTGGAGAAG        TCACCTTCCA        TTTGTCTTAG

4600 TGGTGATTAT   4610 GGGTAGGAAA   4620 ACCTGGTTCT   4630 CCATTCCTGA   4640 GAAGAATCGA
     ACCACTAATA        CCCATCCTTT        TGGACCAAGA        GGTAAGGACT        CTTCTTAGCT
4650 CCTTTAAAGG   4660 ACAGAATTAA   4670 TATAGTTCTC   4680 AGTAGAGAAC
     GGAAATTTCC        TGTCTTAATT        ATATCAAGAG        TCATCTCTTG

4690 TCAAAGAACC   4700 ACCACGAGGA   4710 GCTCATTTTC   4720 TTGCCAAAAG   4730 TTTGGATGAT
     AGTTTCTTGG        TGGTGCTCCT        CGAGTAAAAG        AACGGTTTTC        AAACCTACTA
4740 GCCTTAAGAC   4750 TTATTGAACA   4760 ACCGGAATTG   4770 GCAAGTAAAG
     CGGAATTCTG        AATAACTTGT        TGGCCTTAAC        CGTTCATTTC

4780 TAGACATGGT   4790 TTGGATAGTC   4800 GGAGGCAGTT   4810 CTGTTTACCA   4820 GGAAGCCATG
     ATCTGTACCA        AACCTATCAG        CCTCCGTCAA        GACAAATGGT        CCTTCGGTAC
4830 AATCAACCAG   4840 GCCACCTTAG   4850 ACTCTTTGTG   4860 ACAAGGATCA
     TTAGTTGGTC        CGGTGGAATC        TGAGAAACAC        TGTTCCTAGT

4870 TGCAGGAATT   4880 TGAAAGTGAC   4890 ACGTTTTTCC   4900 CAGAAATTGA   4910 TTTGGGGAAA
     ACGTCCTTAA        ACTTTCACTG        TGCAAAAAGG        GTCTTTAACT        AAACCCCTTT
4920 TATAAACTTC   4930 TCCCAGAATA   4940 CCCAGGCGTC   4950 CTCTCTGAGG
     ATATTTGAAG        AGGGTCTTAT        GGGTCCGCAG        GAGAGACTCC

4960 TCCAGGAGGA   4970 AAAAGGCATC   4980 AAGTATAAGT   4990 TTGAAGTCTA   5000 CGAGAAGAAA
     AGGTCCTCCT        TTTTCCGTAG        TTCATATTCA        AACTTCAGAT        GCTCTTCTTT
5010 GACTAACAGG   5020 AAGATGCTTT   5030 CAAGTTCTCT   5040 GCTCCCCTCC
     CTGATTGTCC        TTCTACGAAA        GTTCAAGAGA        CGAGGGGAGG

5050 TAAAGCTATG   5060 CATTTTTATA   5070 AGACCATGGG   5080 ACTTTTGCTG   5090 GCTTTAGATC
     ATTTCGATAC        GTAAAAATAT        TCTGGTACCC        TGAAAACGAC        CGAAATCTAG
5100 TCTTTGTGAA   5110 GGAACCTTAC   5120 TTTGTGGTTG   5130 TGACATAATT
     AGAAACACTT        CCTTGGAATG        AAGACACCAC        ACTGTATTAA

5140 GGACAAACTA   5150 CCTACAGAGA   5160 TTTAAAGCTC   5170 TAAGGTAAAT   5180 ATAAAATTTT
     CCTGTTTGAT        GGATGTCTCT        AAATTTCGAG        ATTCCATTTA        TATTTTAAAA
5190 TAAGTGTATA   5200 ATGTGTTAAA   5210 CTACTGATTC   5220 TAATTGTTTG
     ATTCACATAT        TACACAATTT        GATGACTAAG        ATTAACAAAC

5230 ATTCCAACCT   5240 ATGGAACTGA   5250 TGAATGGGAG   5260 CAGTGGTGGA   5270 ATGCCTTTAA
     TAAGGTTGGA        TACCTTGACT        ACTTACCCTC        GTCACCACCT        TACGGAAATT
5280 TGAGGAAAAC   5290 CTGTTTTGCT   5300 CAGAAGAAAT   5310 GAGAAAGGTA
     ACTCCTTTTG        GACAAAACGA        GTCTTCTTTA        CTCTTTCCAT

5320 GCCATCTAGT   5330 GATGATGAGG   5340 CTACTGCTGA   5350 CTCTCAACAT   5360 TCTACTCCTC
     CGGTAGATCA        CTACTACTCC        GATGACGACT        GAGAGTTGTA        AGATGAGGAG
5370 CAAAAAAGAA   5380 GAGAAAGGTA   5390 GAAGACCCCA   5400 AGGACTTTCC
     GTTTTTTCTT        CTCTTTCCAT        CTTCTGGGGT        TCCTGAAAGG
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Figure 14F
(SEQ ID NO.: 10 – Primary Sequence)
(SEQ ID NO.: 28 – Complement)

pD17-cJ-dCH2.H1

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5410       5420       5430       5440       5450       5460       5470       5480       5490
TTCAGAATTG CTAAGTTTTT TGAGTCATGC TGTGTTTAGT AATAGAACTC TTGCTTGCTT TGCTATTTAC ACCACAAAGG AAAAAGCTGC
AAGTCTTAAC GATTCAAAAA ACTCAGTACG ACACAAATCA TTATCTTGAG AACGAACGAA ACGATAAATG TGGTGTTTCC TTTTTCGACG

5500       5510       5520       5530       5540       5550       5560       5570       5580
ACTGCTATAC AAGAAAATTA TGGAAAAATA TTCTGTAACC TTTATAAGTA GGCATAACAG TTATAATCAT AACATACTGT TTTTTCTTAC
TGACGATATG TTCTTTTAAT ACCTTTTTAT AAGACATTGG AAATATTCAT CCGTATTGTC AATATTAGTA TTGTATGACA AAAAAGAATG

5590       5600       5610       5620       5630       5640       5650       5660       5670
TCCACACAGG CATAGAGTGT CTGCTATTAA TAACTATGCT CAAAAATTGT GTACCTTTAG CTTTTTAATT TGTAAAGGGG TTAATAAGGA
AGGTGTGTCC GTATCTCACA GACGATAATT ATTGATACGA GTTTTTAACA CATGGAAATC GAAAAATTAA ACATTTCCCC AATTATTCCT

5680       5690       5700       5710       5720       5730       5740       5750       5760
ATATTTGATG TATAGTGCCT TGACTAGAGA TCATAATCAG CCATACCACA TTTGTAGAGG TTTAAAAAAC CTCCCACACC
TATAAACTAC ATATCACGGA ACTGATCTCT AGTATTAGTC GGTATGGTGT AAACATCTCC AAATTTTTTG GAGGGTGTGG

5770       5780       5790       5800       5810       5820       5830       5840       5850
TCCCCCTGAA CCTGAAACAT AAAATGAATG CAATTGTTGT TGTTAACTTG TTTATTGCAG CTTATAATGG TTACAAATAA AGCAATAGCA
AGGGGGACTT GGACTTTGTA TTTTACTTAC GTTAACAACA ACAATTGAAC AAATAACGTC GAATATTACC AATGTTTATT TCGTTATCGT

5860       5870       5880       5890       5900       5910       5920       5930       5940
TCACAAATTT CACAAATAAA GCATTTTTTT CACTGCATTC TAGTGTGTGT TTGTCCAAAC TCATCAATGT ATCTTATCAT GTCTGGATCG
AGTGTTTAAA GTGTTTATTT CGTAAAAAAA GTGACGTAAG ATCACACACA AACAGGTTTG AGTAGTTACA TAGAATAGTA CAGACCTAGC

5950       5960       5970       5980       5990       6000       6010       6020       6030
GCTGGATGAT CCTCCAGCGC GGGGATCTCA TGCTGGAGTT CTTCGCCCAC CCCAACTTGT TTATTGCAGC TTATAATGGT TACAAATAAA
CGACCTACTA GGAGGTCGCG CCCCTAGAGT ACGACCTCAA GAAGCGGGTG GGGTTGAACA AATAACGTCG AATATTACCA ATGTTTATTT

6040       6050       6060       6070       6080       6090       6100       6110       6120
GCAATAGCAT CACAAATTTC ACAAATAAAG CATTTTTTTC ACTGCATTCT AGTTGTGGTT TGTCCAAACT CATCAATGTA TCTTATCATG
CGTTATCGTA GTGTTTAAAG TGTTTATTTC GTAAAAAAAG TGACGTAAGA TCAACACCAA ACAGGTTTGA GTAGTTACAT AGAATAGTAC

6130       6140       6150       6160       6170       6180       6190       6200       6210
TCTGTATACC GTCGACCTCT AGCTAGAGCT ATGGTCATAG ACTGCATTCT CTGTTTCCTG TTGAAATTG  TTATCCGCTC ACAATTCCAC
AGACATATGG CAGCTGGAGA TCGATCTCGA TACCAGTATC TGACGTAAGA GACAAAGGAC GACACTTTAAC AATAGGCGAG TGTTAAGGTG

6220       6230       6240       6250       6260       6270       6280       6290       6300
ACAACATACG AGCCGGAAGC ATAAAGTGTA AAGCCTGGGG TGCCTAATGA GTGAGCTAAC TCACATTAAT TGCGTTGCGC TCACTGCCCG
TGTTGTATGC TCGGCCTTCG TATTTCACAT TTCGGACCCC ACGGATTACT CACTCGATTG AGTGTAATTA ACGCAACGCG AGTGACGGGC
```

Figure 14G
(SEQ ID NO.: 10 – Primary Sequence)
(SEQ ID NO.: 28 – Complement)

pD17-eJ-dCH2.H1

```
       6310        6320        6330        6340        6350        6360        6370        6380        6390
CTTTCCAGTC  GGGAAACCTG  TCGTGCCAGC  TGCATTAATG  AATCGGCCAA  CGGCGGGGGA  GAGGCGGTTT  GCGTATTGGG  CGCTCTTCCG
GAAAGGTCAG  CCCTTTGGAC  AGCACGGTCG  ACGTAATTAC  TTAGCCGGTT  GCGCGCCCCT  CTCCGCCAAA  CGCATAACCC  GCGAGAAGGC

       6400        6410        6420        6430        6440        6450        6460        6470        6480
CTTTCCTGCT  CACTGACTCG  CTGCGCTCGG  TCGTTCGGCT  GCGGCGAGCG  GTATCAGCTC  ACTCAAAGGC  GGTAATACGG  TTATCCACAG
GAAAGGACGA  GTGACTGAGC  GACGCGAGCC  AGCAAGCCGA  CGCCGCTCGC  CATAGTCGAG  TGAGTTTCCG  CCATTATGCC  AATAGGTGTC

       6490        6500        6510        6520        6530        6540        6550        6560        6570
AATCAGGGGA  TAACGCAGGA  AAGAACATGT  GAGCAAAAGG  CCAGCAAAAG  GCCAGGAACC  GTAAAAAGGC  CGCGTTGCTG  GCGTTTTTCC
TTAGTCCCCT  ATTGCGTCCT  TTCTTGTACA  CTCGTTTTCC  GGTCGTTTTC  CGGTCCTTGG  CATTTTTCCG  GCGCAACGAC  CGCAAAAAGG

       6580        6590        6600        6610        6620        6630        6640        6650        6660
ATAGGCTCCG  CCCCCCTGAC  GAGCATCACA  AAAATCGACG  CTCAAGTCAG  AGGTGGCGAA  ACCCGACAGG  ACTATAAAGA  TACCAGGCGT
TATCCGAGGC  GGGGGGACTG  CTCGTAGTGT  TTTTAGCTGC  GAGTTCAGTC  TCCACCGCTT  TGGGCTGTCC  TGATATTTCT  ATGGTCCGCA

       6670        6680        6690        6700        6710        6720        6730        6740        6750
TTCCCCCTGG  AAGCTCCCTC  GTGCGCTCTC  CTGTTCCGAC  CCTGCCGCTT  ACCGGATACC  TGTCCGCCTT  TCTCCCTTCG  GGAAGCGTGG
AAGGGGGACC  TTCGAGGGAG  CACGCGAGAG  GACAAGGCTG  GGACGGCGAA  TGGCCTATGG  ACAGGCGGAA  AGAGGGAAGC  CCTTCGCACC

       6760        6770        6780        6790        6800        6810        6820        6830        6840
CGCTTTCTCA  ATGCTCACGC  TGTAGGTATC  TCAGTTCGGT  GTAGGTCGTT  CGCTCCAAGC  TGGGCTGTGT  GCACGAACCC  CCCGTTCAGC
GCGAAAGAGT  TACGAGTGCG  ACATCCATAG  AGTCAAGCCA  CATCCAGCAA  GCGAGGTTCG  ACCCGACACA  CGTGCTTGGG  GGGCAAGTCG

       6850        6860        6870        6880        6890        6900        6910        6920        6930
CCGACCGCTG  CGCCTTATCC  GGTAACTATC  GTCTTGAGTC  CAACCCGGTA  AGACACGACT  TATCGCCACT  GGCAGCAGCC  ACTGGTAACA
GGCTGGCGAC  GCGGAATAGG  CCATTGATAG  CAGAACTCAG  GTTGGGCCAT  TCTGTGCTGA  ATAGCGGTGA  CCGTCGTCGG  TGACCATTGT

       6940        6950        6960        6970        6980        6990        7000        7010        7020
GGATTAGCAG  AGCGAGGTAT  GTAGGCGGTG  CTACAGAGTT  CTTGAAGTGG  TGGCCTTACT  ACGGCTACAC  TAGAAGGACA  GTATTTGGTA
CCTAATCGTC  TCGCTCCATA  CATCCGCCAC  GATGTCTCAA  GAACTTCACC  ACCGGAATGA  TGCCGATGTG  ATCTTCCTGT  CATAAACCAT

       7030        7040        7050        7060        7070        7080        7090        7100        7110
TCTGCGCTCT  GCTGAAGCCA  GTTACCTTCG  GAAAAAGAGT  TGGTAGCTCT  TGATCCGGCA  AACAAACCAC  CGCTGGTAGC  GGTGGTTTTT
AGACGCGAGA  CGACTTCGGT  CAATGGAAGC  CTTTTTCTCA  ACCATCGAGA  ACTAGGCCGT  TTGTTTGGTG  GCGACCATCG  CCACCAAAAA

       7120        7130        7140        7150        7160        7170        7180        7190        7200
TTGTTTGCAA  GCAGCAGATT  ACCGCGAGAA  AAAAAGGATC  TCAAGAAGAT  CCTTTGATCT  TTTCTACGGG  GTCTGACGCT  CAGTGGAACG
AACAAACGTT  CGTCGTCTAA  TGGCGCTCTT  TTTTTCCTAG  AGTTCTTCTA  GGAAACTAGA  AAAGATGCCC  CAGACTGCGA  GTCACCTTGC
```

Figure 14H
(SEQ ID NO.: 10 – Primary Sequence)
(SEQ ID NO.: 28 – Complement)

pD17-cJ-dCH2.H1

```
7210        7220        7230        7240        7250        7260        7270        7280        7290
AAAACTCACG  TTAAGGGATT  TTGGTCATGA  GATTATCAAA  AAGGATCTTC  ACCTAGATCC  TTTTAAATTA  AAAATGAAGT  TTTAAATCAA
TTTTGAGTGC  AATTCCCTAA  AACCAGTACT  CTAATAGTTT  TTCCTAGAAG  TGGATCTAGG  AAAATTTAAT  TTTTACTTCA  AAATTTAGTT

7300        7310        7320        7330        7340        7350        7360        7370        7380
TCTAAAGTAT  ATATGAGTAA  ACTTGGTCTG  ACAGTTACCA  ATGCTTAATC  AGTGAGGCAC  CTATCTCAGC  GATCGTCTTA  TTTCGTTCAT
AGATTTCATA  TATACTCATT  TGAACCAGAC  TGTCAATGGT  TACGAATTAG  TCACTCCGTG  GATAGAGTCG  CTAGACAGAT  AAAGCAAGTA

7390        7400        7410        7420        7430        7440        7450        7460        7470
CCATAGTTGC  CTGACTCCCC  GTCGTGTAGA  TAACTACGAT  ACGGGAGGGC  TTACCATCTG  GCCCCAGTGC  TGCAATGATA  CCGGCGAGACC
GGTATCAACG  GACTGAGGGG  CAGCACATCT  ATTGATGCTA  TGCCCTCCCG  AATGGTAGAC  CGGGGTCACG  ACGTTACTAT  GGCGGCTCTGG

7480        7490        7500        7510        7520        7530        7540        7550        7560
CACGCTCACC  GGCTCCAGAT  TTATCAGCAA  TAAACCAGCC  AGCCGGAAGG  GCCGAGCGCA  GAAGTGGTCC  TGCAACTTTA  TCCGCCTCCA
GTGCGAGTGG  CCGAGGTCTA  AATAGTCGTT  ATTTGGTCGG  TCGGCCTTCC  CGGCTCGCGT  CTTCACCAGG  ACGTTGAAAT  AGGCGGAGGT

7570        7580        7590        7600        7610        7620        7630        7640        7650
TCCAGTCTAT  TAATTGTTGC  CGGGAAGCTA  GAGTAAGTAG  TTCGCCAGTT  AATAGTTTGC  GCAACGTTGT  TGCCATTGCT  ACAGGCATCG
AGGTCAGATA  ATTAACAACG  GCCCTTCGAT  CTCATTCATC  AAGCGGTCAA  TTATCAAACG  CGTTGCAACA  ACGGTAACGA  TGTCCGTAGC

7660        7670        7680        7690        7700        7710        7720        7730        7740
TGGTGTCACG  CTCGTCGTTT  GGTATGGCTT  CATTCAGCTC  CGGTTCCCAA  CGATCAAGGC  GAGTTACATG  ATCCCCCATG  TTGTGCAAAA
ACCACAGTGC  GAGCAGCAAA  CCATACCGAA  GTAAGTCGAG  GCCAAGGGTT  GCTAGTTCCG  CTCAATGTAC  TAGGGGGTAC  AACACGTTTT

7750        7760        7770        7780        7790        7800        7810        7820        7830
AAGCGGTTAG  CTCCTTCGGT  CCTCCGATCG  TTGTCAGAAG  TAAGTTGGCC  GCAGTGTTAT  CACTCATGGT  TATGGCAGCA  CTGCATAATT
TTCGCCAATC  GAGGAAGCCA  GGAGGCTAGC  AACAGTCTTC  ATTCAACCGG  CGTCACAATA  GTGAGTACCA  ATACCGTCGT  GACGTATTAA

7840        7850        7860        7870        7880        7890        7900        7910        7920
CTCTTACTGT  CATGCCATCC  GTAAGATGCT  TTTCTGTGAC  TGGTGAGTAC  TCAACCAAGT  CATTCTCGAGA ATAGTGTATG  CGGGGACCGA
GAGAATGACA  GTACGGTAGG  CATTCTACGA  AAAGACACTG  ACCACTCATG  AGTTGGTTCA  GTAAGACTCT  TATCACATAC  GCCCCTGGCT

7930        7940        7950        7960        7970        7980        7990        8000        8010
GTTGCTCTTG  CCCGGCGTCA  ATACGGGATA  ATACCGCGCC  ACATAGCAGA  ACTTTAAAAG  TGCTCATCAT  TGGAAAACGT  TCTTCGGGGC
CAACGAGAAC  GGGCCGCAGT  TATGCCCTAT  TATGGCGCGG  TGTATCGTCT  TGAAATTTTC  ACGAGTAGTA  ACCTTTTGCA  AGAAGCCCCG

8020        8030        8040        8050        8060        8070        8080        8090        8100
GAAAACTCTC  AAGGATCGTTA CCGCTCGTTGA GATCCGAGTTC GATGTAACCC  ACTCGTGCAC  CCAACTGATC  TTCAGCATCT  TTTACTTTCA
CTTTTGAGAG  TTCCTAGCAAT GGCGACAACT  CTAGGTCAAG  CTACATTGGG  TGAGCACGTG  GGTTGACTAG  AAGTCGTAGA  AAATGAAAGT
```

Figure 14I
(SEQ ID NO.: 10 – Primary Sequence)
(SEQ ID NO.: 28 – Complement)

pD17-cJ-dCH2.H1

```
        8110        8120        8130        8140        8150        8160        8170        8180        8190
  CCAGCGTTTC  TGGGTGAGCA  AAAACAGGAA  GGCAAATGC   CGCAAAAAG   GGAATAAGGG  CGACACGGAA  ATGTTGAATA  CTCATACTCT
  GGTCGCAAAG  ACCCACTCGT  TTTTGTCCTT  CCGTTTACG   GCGTTTTTC   CCTTATTCCC  GCTGTGCCTT  TACAACTTAT  GAGTATGAGA

        8200        8210        8220        8230        8240        8250        8260        8270        8280
  TCCTTTTTCA  ATATTATTGA  AGCATTTATC  AGGGTTATTG  TCTTCATGAGC GGATACATAT  TTGAATGTAT  TTAGAAAAAT  AAACAAATAG
  AGGAAAAAGT  TATAATAACT  TCGTAAATAG  TCCCAATAAC  AGAGTACTCG  CCTATGTATA  AACTTACATA  AATCTTTTTA  TTTGTTTATC

        8290        8300        8310        8320        8330
  GGGTTCCGCG  CACATTTCCC  CGAAAGTGC   CACCTGACGT  C
  CCCAAGGCGC  GTGTAAAGGG  GCTTTTCACG  GTGGACTGCA  G
```

Figure 14J
(SEQ ID NO.: 10 – Primary Sequence)
(SEQ ID NO.: 28 – Complement)

Figure 19A
(SEQ ID NO.: 23 – Primary Sequence)
(SEQ ID NO.: 29 – Complement)

pD17-hG1b

```
        10         20         30         40         50         60
GGTACCAATT TAAATTGATA TCTCCTTAGG TCTCGAGTCT CTAGATAACC GGTCAATCGA
CCATGGTTAA ATTTAACTAT AGAGGAATCC AGAGCTCAGA GATCTATTGG CCAGTTAGCT

        70         80         90        100        110        120
TTGGAATTCT TGCGGCCGCT TGCTAGCACC AAGGGCCCAT CGGTCTTCCC CCTGGCACCC
AACCTTAAGA ACGCCGGCGA ACGATCGTGG TTCCCGGGTA GCCAGAAGGG GGACCGTGGG

       130        140        150        160        170        180
TCCTCCAAGA GCACCTCTGG GGGCACAGCG GCCCTGGGCT GCCTGGTCAA GGACTACTTC
AGGAGGTTCT CGTGGAGACC CCCGTGTCGC CGGGACCCGA CGGACCAGTT CCTGATGAAG

       190        200        210        220        230        240
CCCGAACCGG TGACGGTGTC GTGGAACTCA GGCGCCCTGA CCAGCGGCGT GCACACCTTC
GGGCTTGGCC ACTGCCACAG CACCTTGAGT CCGCGGGACT GGTCGCCGCA CGTGTGGAAG

       250        260        270        280        290        300
CCGGCTGTCC TACAGTCCTC AGGACTCTAC TCCCTCAGCA GCGTGGTCAC CGTGCCCTCC
GGCCGACAGG ATGTCAGGAG TCCTGAGATG AGGGAGTCGT CGCACCAGTG GCACGGGAGG

       310        320        330        340        350        360
AGCAGCTTGG GCACCCAGAC CTACATCTGC AACGTGAATC ACAAGCCCAG CAACACCAAG
TCGTCGAACC CGTGGGTCTG GATGTAGACG TTGCACTTAG TGTTCGGGTC GTTGTGGTTC

       370        380        390        400        410        420
GTGGACAAGA AAGTTGGTGA GAGGCCAGCA CAGGGAGGGA GGGTGTCTGC TGGAAGCCAG
CACCTGTTCT TTCAACCACT CTCCGGTCGT GTCCCTCCCT CCCACAGACG ACCTTCGGTC

       430        440        450        460        470        480
GCTCAGCGCT CCTGCCTGGA CGCATCCCGG CTATGCAGCC CCAGTCCAGG GCAGCAAGGC
CGAGTCGCGA GGACGGACCT GCGTAGGGCC GATACGTCGG GGTCAGGTCC CGTCGTTCCG

       490        500        510        520        530        540
AGGCCCCGTC TGCCTCTTCA CCCGGGAGGCC TCTGCCCGCC CCACTCATGC TCAGGGAGAG
TCCGGGGCAG ACGGAGAAGT GGGCCTCCGG AGACGGGCGG GGTGAGTACG AGTCCCTCTC

       550        560        570        580        590        600
GGTCTTCTGG CTTTTTCCCC AGGCACACGG CTAGGTGCCC TCAAGCCAGG
CCAGAGACC GAAAAAGGGG TCCGAGACCC GATCCACGGG GATTGGGTCC
```

Figure 19B
(SEQ ID NO.: 23 – Primary Sequence)
(SEQ ID NO.: 29 – Complement)

pD17-hG1ab

```
        610        620        630        640        650        660
CCCTGCACAC AAAGGGGCAG GTGCTGGGCT CAGACCTGCC AAGAGCCATA TCCGGGGAGGA
GGGACGTGTG TTTCCCCGTC CACGACCCGA GTCTGGACGG TTCTCGGTAT AGGCCCCTCCT

        670        680        690        700        710        720
CCCTGCCCCT GACCTAAGCC CACCCCAAAG GCCAAAACTCT CCACTCCCTC AGCTCGGACA
GGGACGGGGA CTGGATTCGG GTGGGGTTTC CGGTTTGAGA GGTGAGGGAG TCGAGCCTGT

        730        740        750        760        770        780
CCTTCTCTCC TCCCAGATTC CAGTAACTCC CAATCTTCTC TCTGCAGAGC CCAAATCTTG
GGAAGAGAGG AGGGTCTAAG GTCATTGAGG GTTAGAAGAG AGACGTCTCG GGTTTAGAAC

        790        800        810        820        830        840
TGACAAAACT CACACATGCC CACCGTGCCC AGGTAAGCCA GCCCAGGCCT CGCCCTCCAG
ACTGTTTTGA GTGTGTACGG GTGGCACGGG TCCATTCGGT CGGGTCCGGA GCGGGAGGTC

        850        860        870        880        890        900
CTCAAGGCGG GACAGGTGCC CTAGAGTAGC CTGCATCCAG GGACAGGCGCC CAGCGCGGGTG
GAGTTCCGCC CTGTCCACGG GATCTCATCG GACGTAGGTC CCTGTCCGGG GTCGGGCCCAC

        910        920        930        940    950 235      960
CTGACACGTC CACCTCCATC TCTTCCTCAG CACCTGAACT CCTGGGGGGA CCGTCAGTCT
GACTGTGCAG GTGGAGGTAG AGAAGGAGTC GTGGACTTGA 237 GGACCCCCCT GGCAGTCAGA

        970        980        990        1000       1010       1020
TCCTCTTCCC CCCAAAACCC AAGGACACCC TCATGATCTC CCGGACCCCT GAGGTCACAT
AGGAGAAGGG GGGTTTTGGG TTCCTGTGGG AGTACTAGAG GGCCTGGGGA CTCCAGTGTA

        1030       1040       1050       1060       1070       1080
GCGGTGGTGCT GGACGTGAGC CACGAAGACC CTGAGGTCAA GTTCAACTGG TACGTGGACG
CGCACCACCA CCTGCACTCG GTGCTTCTGG GACTCCAGTT CAAGTTGACC ATGCACCTGC

        1090       1100       1110       1120       1130       1140
GCGTGGAGGT GCATAATGCC AAGACAAAGC CGCGGGAGGA GCAGTACAAC AGCACGTACC
CGCACCTCCA CGTATTACGG TTCTGTTTCG GCGCCCTCCT CGTCATGTTG TCGTGCATGG

        1150       1160       1170    1180  1190 318   1200 320
CTGTGGTCAG CGTCCTGCAC AGGACTGGCT GAATGGCAAG GAGTACAAGT
CACACCAGTC GCAGGACGTG TCCTGACCGA CTTACCGTTC CTCATGTTCA
```

Figure 19C
(SEQ ID NO.: 23 – Primary Sequence)
(SEQ ID NO.: 29 – Complement)

pD17-hG1b

```
      1210       1220       1230       1240       1250       1260
GCAAGGTCTC CAACAAAGCC CTCCCAGCCC CCATCGAGAA AACCATCTCC AAAGCCAAAG
CGTTCCAGAG GTTGTTTCGG GAGGGTCGGG GGTAGCTCTT TTGGTAGAGG TTTCGGTTTC

      1270       1280       1290       1300       1310       1320
GTGGGGACCCG TGGGGTGCGA GGGCCCACATG GACAGAGGCC GGCTCGGCCC ACCCTCTGCC
CACCCTGGGC ACCCCACGCT CCCGGTGTAC CTGTCTCCGG CCGAGCCGGG TGGGAGACGG

      1330       1340       1350       1360       1370       1380
CTGAGAGTGA CCGGCTGTACC AACCTCTGTC AGCCCCGAGA ACCACAGGTG
GACTCTCACT GGCGACATGG TTGGAGACAG GGATGTCCCG TCGGGGCTCT TGGTGTCCAC

      1390       1400       1410       1420       1430       1440
TACACCCTGC CCCCATCCCG ACCAAGAACC AGGTCAGCCT GACCTGCCTG
ATGTGGGACG GGGGTAGGGC CCTACTCGAC TGGTTCTTGG CTGGACGGAC

      1450       1460       1470       1480       1490       1500
GTCAAAGGCT TCTATCCCAG CGACATCGCC GTGGAGTGGG AGAGCAATGG GCAGCCGGAG
CAGTTTCCGA AGATAGGGTC GCTGTAGCGG CACCTCACCC TCTCGTTACC CGTCGGCCTC

      1510       1520       1530       1540       1550       1560
AACAACTACA AGACCACGCC TCCCGTGCTG GACTCCGACG GCTCCTTCTT CCTCTACAGC
TTGTTGATGT TCTGGTGCGG AGGGCACGAC CTGAGGCTGC CGAGGAAGAA GGAGATGTCG

      1570       1580       1590       1600       1610       1620
AAGCTCACCG TGGACAAGAG CAGGTGGCAG CAGGGGAACG TCTTCTCATG CTCCGTGATG
TTCGAGTGGC ACCTGTTCTC GTCCACCGTC GTCCCCTTGC AGAAGAGTAC GAGGCACTAC

      1630       1640       1650       1660       1670       1680
CATGAGGCTC TGCACAACCA CTACACGCAG AAGAGCCTCT CCCTGTCTCC GGGTAAATGA
GTACTCCGAG ACGTGTTGGT GATGTGCGTC TTCTCGGAGA GGGACAGAGG CCCATTTACT

      1690       1700       1710       1720       1730       1740
GTGGCGACGGC CGGCTCCCCG GGCTCTCGCG GTCGGCACGAG GATGCTTGGC
CACCGCTGCCG GCCGAGGGGC CCGAGAGCGC CAGCGTGCTC CTACGAACCG

      1750       1760       1770       1780       1790       1800
ACGTACCCCC TGTTACATACT TCCCGGGGGC CCAGCCATGGA GTCGGCACTGC
TGCATGGGGG ACAATGTATGA AGGGCCCCCG GGTCGGTACCT TTATTTCGTTG GGTCGCGACG
```

Figure 19D
(SEQ ID NO.: 23 – Primary Sequence)
(SEQ ID NO.: 29 – Complement)

pD17-hG1b

```
     1810       1820       1830       1840       1850       1860
CCTGGGCCCC TGCGAGACTG TGATGGTTCT TTCCACGGGT CAGGCCGAGT CTGAGGCCTG
GGACCCGGGG ACGCTCTGAC ACTACCAAGA AAGGTGCCCA GTCCGGCTCA GACTCCGGAC

     1870       1880       1890       1900       1910       1920
AGTGGCATGA GGGAGGCAGA GCGGGTCCCA CTGTCCCCAC ACTGGCCCAG GCTGTGCAGG
TCACCGTACT CCCTCCGTCT CGCCCAGGGT GACAGGGGTG TGACCGGGTC CGACACGTCC

     1930       1940       1950       1960       1970       1980
TGTGCCTGGG CCCCCTAGGG TGGGGCTCAG CCAGGGCTG  CCCTCGGCAG GGTGGGGGAT
ACACGGACCC GGGGGATCCC ACCCGAGTC  GGTCCCGAC  GGGAGCCGTC CCACCCCCTA

     1990       2000       2010       2020       2030       2040
TTGGCCAGCGT GGCCCTCCCT CCAGCAGCAC CTGCCCTGGG CTGGGCCACG GGAAGCCCTA
AACGGTCGCA  CCGGGAGGGA GGTCGTCGTG GACGGGACCC GACCCGGTGC CCTTCGGGAT

     2050       2060       2070       2080       2090       2100
GGAGCCCCTG GGGACAGACA CACAGCCCCT GCCTCTGTAG GAGACTGTCC TGTTCTGTGA
CCTCGGGGAC CCCTGTCTGT GTGTCGGGGA CGGAGACATC CTCTGACAGG ACAAGACACT

     2110       2120       2130       2140       2150       2160
GCGCCCCTGT CCTCCCCGACC TCCATGCCCA CTCGGGGGCA TGCTGGGGAT GCGGGTGGGCT
CGCGGGGACA GGAGGGCTGG  AGGTACGGGT GAGCCCCCGT ACGACCCCTA CGCCCACCGA

     2170       2180       2190       2200       2210       2220
CTATGGCTTC TGAGGCGGAA AGAACCAGCT GGGGCTGTAG GGGGTTATCC CACGCGCCCT
GATACCGAAG ACTCCGCCTT TCTTGGTCGA CCCCGAGATC CCCCATAGGG GTGCGCGGGA

     2230       2240       2250       2260       2270       2280
GTAGCGGCGC ATTAAGCGCG GCGGGTGTGG TGGTTACGCG CAGCGGTGACC GCTACACTTG
CATCGCCGCG TAATTCGCGC CGCCCACACC ACCAATGCGC GTCGCCACTGG CGATGTGAAC

     2290       2300       2310       2320       2330       2340
CCAGCGCCCT AGCGCCCGCT CCTTTCGCTT TCTTCCCTTC CAGCGGTGACC ACGTTCGCCG
GGTCGCGGGA TCGCGGGCGA GGAAAGCGAA AGAAGGGAAG GTCGCCACTG  TGCAAGCGGC

     2350       2360       2370       2380       2390       2400
GCTTTCCCCG TCAAGCTCTA AATCGGGGCA TCCCTTTAGG GTTCCGATTT AGTGCTTTAC
CGAAAGGGGC AGTTCGAGAT TTAGCCCCGT AGGGAAATCC CAAGGCTAAA TCACGAAATG
```

Figure 19E
(SEQ ID NO.: 23 – Primary Sequence)
(SEQ ID NO.: 29 – Complement)

pD17-hG1b

```
       2410        2420        2430        2440        2450        2460
   GGCACCTCGA  CCCCAAAAAA  CTTGATTAGG  GTGATGGTTC  ACGTAGTGGG  CCATCGCCCT
   CCGTGGAGCT  GGGGTTTTTT  GAACTAATCC  CACTACCAAG  TGCATCACCC  GGTAGCGGGA

       2470        2480        2490        2500        2510        2520
   GATAGACGGT  TTTTCGCCCT  TTGACGTTGG  AGTCCACGTT  CTTTAATAGT  GGACTCTTGT
   CTATCTGCCA  AAAAGCGGGA  AACTGCAACC  TCAGGTGCAA  GAAATTATCA  CCTGAGAACA

       2530        2540        2550        2560        2570        2580
   TCCAAACTGG  AACAACACTC  CGGTCTATTC  TTTTGATTTA  TTTTGATTTA  TAAGGGATTT
   AGGTTTGACC  TTGTTGTGAG  GCCAGATAAG  AAAACTAAAT  AAAACTAAAT  ATTCCCTAAA

       2590        2600        2610        2620        2630        2640
   TGGGGATTTC  GGCCTATTGG  TTAAAAAATG  AGCTGATTTA  ACAAAAATTT  AACGGCGAATT
   ACCCCTAAAG  CCGGATAACC  AATTTTTTAC  TCGACTAAAT  TGTTTTTAAA  TTGCGCTTAA

       2650        2660        2670        2680        2690        2700
   AATTCTGTGG  AGTTAGGGTG  TGGAAAGTCC  CCAGGCTCCC  CAGGCAGGCA  ...
   TTAAGACACC  TCAATCCCAC  ACCTTTCAGG  GGTCCGAGGG  GTCCGTCCGT

       2710        2720        2730        2740        2750        2760
   GAAGTATGCA  AAGCATGCAT  CTCAAATAGT  CAGCAACCAT  AGTCCCGCCC  CTAACTCCGC
   CTTCATACGT  TTCGTACGTA  GAGTTTATCA  GTCGTTGGTA  TCAGGGCGGG  GATTGAGGCG

       2770        2780        2790        2800        2810        2820
   CCATCCCGCC  CCTAACTCCG  CCCAGTTCCG  CCCATTCTCC  GCCCCATGGC  TTGACTAATTT
   GGTAGGGCGG  GGATTGAGGC  GGGTCAAGGC  CGGGGTACCG  ACTGATTAAA

       2830        2840        2850        2860        2870        2880
   TTTTTATTTA  TGCAGAGGCC  GAGGCCGCCT  CGGCCTCTGA  GCTATTCCAG  AAGTAGTGAG
   AAAAATAAAT  ACGTCTCCGG  CTCCGGCGGA  GCCGGAGACT  CGATAAGGTC  TTCATCACTC

       2890        2900        2910        2920        2930        2940
   GAGGCTTTTT  TGGAGGGCTA  GGCTTTTGCA  AAAAGCTTGG  ACAGCTCAGG  GCTGCGGATTT
   CTCCGAAAAA  ACCTCCGGAT  CCGAAAACGT  TTTTCGAACC  TGTCGAGTCC  CGACGGCTAAA

       2950        2960        2970        2980        2990        3000
   CGGGCCAAAC  TTGACGGGCAA  TCCTAGCGGTG  AAGGCTGGTA  GGATTTTATC  CCCGGCTGCCA
   GCGGGGTTTG  AACTGCCGTT  AGGATCGCAC  TTCCGACCAT  CCTAAAATAG  GGGCGGACGGT
```

Figure 19F
(SEQ ID NO.: 23 – Primary Sequence)
(SEQ ID NO.: 29 – Complement)

pD17-hG1b

```
3010         3020         3030         3040         3050         3060
TCATGGTTCG   ACCATTGAAC   TGCATCGTCG   CCGTGTCCCA   AAATATGGGG   ATTGGCAAGA
AGTACCAAGC   TGGTAACTTG   ACGTAGCAGC   GGCACAGGGT   TTTATACCCC   TAACCGTTCT

3070         3080         3090         3100         3110         3120
ACGGAGACCT   CCCTGGCCT    CCGCTCAGGA   ACGAGTTCAA   GTACTTCCAA   AGAATGACCA
TGCCTCTGGA   TGGGACCGGA   GGCGAGTCCT   TGCTCAAGTT   CATGAAGGTT   TCTTACTGGT

3130         3140         3150         3160         3170         3180
CAACCTCTTC   AGTGGAAGGT   AAACAGAATC   TGGTGATTAT   GGGTAGGAAA   ACCTGGTTCT
GTTGGAGAAG   TCACCTTCCA   TTTGTCTTAG   ACCACTAATA   CCCATCCTTT   TGGACCAAGA

3190         3200         3210         3220         3230         3240
CCATTCCTGA   GAAGAATCGA   CCTTTAAAGG   ACAGAATTAA   TATAGTTCTC   AGTAGAGAAC
GGTAAGGACT   CTTCTTAGCT   GGAAATTCC   TGTCTTAATT   ATATCAAGAG   TCATCTCTTG

3250         3260         3270         3280         3290         3300
TCAAAGAACC   ACCACGAGGA   GCTCATTTTC   TTGCCAAAAG   TTTGGATGAT   GCCTTAAGAC
AGTTTCTTGG   TGGTGCTCCT   CGAGTAAAAG   AACGGTTTTC   AAACCTACTA   CGGAATTCTG

3310         3320         3330         3340         3350         3360
TTATTGAACA   ACCGGAATTG   GCAAGTAAAG   TAGACATGGT   TTGGATAGTC   GGAGGCAGTT
AATAACTTGT   TGGCCTTAAC   CGTTCATTTC   ATCTGTACCA   AACCTATCAG   CCTCCGTCAA

3370         3380         3390         3400         3410         3420
CTGTTTACCA   GGAAGCCATG   AATCAACCAG   GCCACCTTAG   ACTCTTTGTG   ACAAGGATCA
GACAAATGGT   CCTTCGGTAC   TTAGTTGGTC   CGGTGGAATC   TGAGAAACAC   TGTTCCTAGT

3430         3440         3450         3460         3470         3480
TGCAGGAATT   TGAAAGTGAC   ACGTTTTTCC   CAGAAATTGA   TTTGGGGAAA   TATAAACTTC
ACGTCCTTAA   ACTTTCACTG   TGCAAAAAGG   GTCTTTAACT   AAACCCCTTT   ATATTTGAAG

3490         3500         3510         3520         3530         3540
TCCCAGAATA   CCCAGGCGTC   CTCTCTGAGG   TCCAGGAGGA   AAAAGGCATC   AAGTATAAGT
AGGGTCTTAT   GGGTCCGCAG   GAGAGACTCC   AGGTCCTCCT   TTTTCCGTAG   TTCATATTCA

3550         3560         3570         3580         3590         3600
TTGAAGTCTA   CGAGAAGAAA   GACTAACAGG   AAGATGCTTT   CAAGTTCTCT   GCTCCCCTCC
AACTTCAGNT   GCTCTTCTTT   CTGATTGTCC   TTCTACGAAA   GTTCAAGAGA   CGAGGGGAGG
```

Figure 19G
(SEQ ID NO.: 23 – Primary Sequence)
(SEQ ID NO.: 29 – Complement)

pD17-hG1b

```
        3610        3620        3630        3640        3650        3660
    TAAAGCTATG  CATTTTTATA  AGACCATGGG  ACTTTTGCTG  GCTTTAGATC  TCTTTGTGAA
    ATTTCGATAC  GTAAAAATAT  TCTGGTACCC  TGAAAACGAC  CGAAATCTAG  AGAAACACTT

        3670        3680        3690        3700        3710        3720
    GGAACCTTAC  TTCTGTGGTG  TGACATAATT  GGACAAACTA  CCTACAGAGA  TTTAAAGCTC
    CCTTGGAATG  AAGACACCAC  ACTGTATTAA  CCTGTTTGAT  GGATGTCTCT  AAATTTCGAG

        3730        3740        3750        3760        3770        3780
    TAAGGTAAAT  ATAAAAATTT  TAAGTGTATA  ATGTGTTAAA  CTACTGATTC  TTAATTGTTTG
    ATTCCATTTA  TATTTTAAAA  ATTCACATAT  TACACAATTT  GATGACTAAG  AATTAACAAAC

        3790        3800        3810        3820        3830        3840
    TGTATTTTAG  ATTCCAACCT  ATGGAACTGA  TGAATGGGAG  CAGTGGTGGA  ATGCCTTTAA
    ACATAAAATC  TAAGGTTGGA  TACCTTGACT  ACTTACCCTC  GTCACCACCT  TACGGAAATT

        3850        3860        3870        3880        3890        3900
    TGAGGAAAAC  CTGTTTTGCT  CAGAAGAAAT  GCCATCTAGT  GATGGTGAGG  CTACTGCTGA
    ACTCCTTTTG  GACAAAACGA  GTCTTCTTTA  CGGTAGATCA  CTACTACTCC  GATGACGACT

        3910        3920        3930        3940        3950        3960
    CTCTCAACAT  TCTACTCCTC  CAAAAAAGAA  GAGAAAGGTA  GAAGACCCCA  AGGACTTTCC
    GAGAGTTGTA  AGATGAGGAG  GTTTTTTCTT  CTCTTTCCAT  CTTCTGGGGT  TCCTGAAAGG

        3970        3980        3990        4000        4010        4020
    TTCAGAATTG  CTAAGTTTTT  TGAGTCATGC  TGTGTTTAGT  AATAGAAGCTC  TTGCTTGCTT
    AAGTCTTAAC  GATTCAAAAA  ACTCAGTACG  ACACAAATCA  TTATCTTCGAG  AACGAACGAA

        4030        4040        4050        4060        4070        4080
    TGCTATTTAC  ACCACAAAGG  AAAAAAGCTGC  ACTGCTATAC  AAGAAAAATTA  TGGAAAAAATA
    ACGATAAATG  TGGTGTTTCC  TTTTTTCGACG  TGACGATATG  TTCTTTTTAAT  ACCTTTTTAT

        4090        4100        4110        4120        4130        4140
    TTCTGTAACC  TTTTATAAGTA  GGCATAAACAG  TTATAATCAT  AACATACTGT  TTTTTTCTTAC
    AAGACATTGG  AAATATTCAT  CCGTATTGTC  AATATTAGTA  TTGTATGACA  AAAAAGAATG

        4150        4160        4170        4180        4190        4200
    TCCACACAGG  CATAGAGTGT  CTGCTATTAA  TAACTATTGCT  CAAAAATTGT  GTACCTTTAG
    AGGTGTGTCC  GTATCTCACA  GACGATAATT  ATTGATATCGA  GTTTTTAACA  CATGGAAATC
```

Figure 19H
(SEQ ID NO.: 23 – Primary Sequence)
(SEQ ID NO.: 29 – Complement)

pD17-hG1b

```
                                4210       4220       4230       4240       4250       4260
CTTTTTAATT TGTAAAGGGG TTAATAAGGA ATATTTGATG TATAGTGCCT TGACTAGAGA
GAAAAATTAA ACAATTCCCC AATTATTCCT TATAAACTAC ATATCACGGA ACTGATCTCT

                                4270       4280       4290       4300       4310       4320
TCATAATCAG CCATACCACA TTTGTAGAGG TTTTTACTTGC TTTAAAAAAC CTCCCACACC
AGTATTAGTC GGTATGGTGT AAACATCTCC AAAATGAACG AAATTTTTTG GAGGGTGTGG

                                4330       4340       4350       4360       4370       4380
TCCCCCCTGAA CCTGAAACAT AAAATGAATG CAATTGTTGT TGTTAACTTG TTTATTGCAG
AGGGGGACTT GGACTTTGTA TTTTACTTAC GTTAACAACA ACAATTGAAC AAATAACGTC

                                4390       4400       4410       4420       4430       4440
CTTATAAATGG TTACAAATAA AGCAAATAGCA TCACAAATTT CACAAAATAAA GCATTTTTTT
GAATATTACC AATGTTTATT TCGTTATCGT AGTGTTTAAA GTGTTTTATTT CGTAAAAAAA

                                4450       4460       4470       4480       4490       4500
CACTGCATTC TAGTTGTTGGT TTGTCCAAAC TCATCAATGT ATCTTTATTCAT GTCTGGATCG
GTGACGTAAG ATCAACACCA AACAGGTTTG AGTAGTTACA TAGAATAGTA CAGACCTAGC

                                4510       4520       4530       4540       4550       4560
GCTGGATGAT CCTCCAGCGC GGGGATCTCA TGCTGGAGTT CTTCGCCCAC CCCAACTTGT
CGACCTACTA GGAGGTCGCG CCCCTAGAGT ACGACCTCAA GAAGCGGGTG GGGTTGAACA

                                4570       4580       4590       4600       4610       4620
TTATTGCAGC TTATAAATGGT TACAAATATAA GCAATAGCAT CACAAATTTC ACAAATAAAG
AATAACGTCG AATATTACCA ATGTTTATTT CGTTATCGTA GTGTTTAAAG TGTTTATTTC

                                4630       4640       4650       4660       4670       4680
CATTTTTTTC ACTGCATTCT AGTTGTGGTT TGTCCAAACT CATCAATGTA TCTTTATCATG
GTAAAAAAAG TGACGTAAGA TCAACACCAA ACAGGTTTGA GTAGTTACAT AGAATAGTAC

                                4690       4700       4710       4720       4730       4740
TCTGTATACC GTCGACCTCT AGCTAGAGCT TGGCGTAATC ATGGTCATAG CTGTTTCCTG
AGACATATGG CAGCTGGAGA TCGATCTCGA ACCGCATTAG TACCAGTATC GACAAAGGAC

                                4750       4760       4770       4780       4790       4800
TGTGAAATTG TTATCCGCTC ACAATTCCAC ACAACATACG AGCCGGAAGC ATAAAGTGTA
ACACTTTAAC AATAGGCGAG TGTTGTATGC TCGGCCTTCG TATTTCACAT
```

Figure 19I
(SEQ ID NO.: 23 – Primary Sequence)
(SEQ ID NO.: 29 – Complement)

pD17-hG1b

```
4810        4820        4830        4840        4850        4860
AAGCCTGGGG  TGCCTAATGA  GTGAGCTAAC  TCACATTAAT  TGCGTTGCGC  TCACTGCCCG
TTCGGACCCC  ACGGATTACT  CACTCGATTG  AGTGTAATTA  ACGCAACGCG  AGTGACGGGC

4870        4880        4890        4900        4910        4920
CTTTTCCAGTC GGGAAACCTG  TCGTGCCAGC  TGCATTAATG  AATCGGCCAA  CGCGCGGGGA
GAAAGGTCAG  CCCTTTGGAC  AGCACGGTCG  ACGTAATTAC  TTAGCCGGTT  GCGCGCCCCT

4930        4940        4950        4960        4970        4980
GAGGCGGTTT  GCGTATTGGG  CGCTCCTTCCG CTTCCTCGCT  CACTGACTCG  CTGCGCTCGG
CTCCGCCAAA  CGCATAACCC  GCGAGGAAGC  GAAGGAGCGA  GTGACTGAGC  GACGCGAGCC

4990        5000        5010        5020        5030        5040
TCGTTCGGCT  GCGGCGGAGCG GTATCAGCTC  ACTCAAAGGC  GGTAATACGG  TTATCCACAG
AGCAAGCCGA  CGCCGCTCGC  CATAGTCGAG  TGAGTTTCCG  CCATTATGCC  AATAGGTGTC

5050        5060        5070        5080        5090        5100
AATCAGGGGA  TAACGCAGGA  AAGAACATGT  GAGCAAAAGG  CCAGCAAAAG  GCCAGGAACC
TTAGTCCCCT  ATTGCGTCCT  TTCTTGTACA  CTCGTTTTCC  GGTCGTTTTC  CGGTCCTTGG

5110        5120        5130        5140        5150        5160
GTAAAAAGGC  CGCGTTGCTG  GCGTTTTTCC  ATAGGCTCCG  CCCCCCTGAC  GAGCATCACA
CATTTTTCCG  GCGCAACGAC  CGCAAAAAGG  TATCCGAGGC  GGGGGGACTG  CTCGTAGTGT

5170        5180        5190        5200        5210        5220
AAAATCGACG  CTCAAGTCAG  AGGTGGCGAA  ACCCGACAGG  ACTATAAAGA  TACCAGGCGT
TTTTAGCTGC  GAGTTCAGTC  TCCACCGCTT  TGGGCTGTCC  TGATATTTCT  ATGGTCCGCA

5230        5240        5250        5260        5270        5280
TTCCCCCTGG  AAGCTCCCTC  GTGCGCTCTC  CTGTTCCGAC  CCTGCCGCTT  ACCGGATACC
AAGGGGGACC  TTCGAGGGAG  CACGCGAGAG  GACAAGGCTG  GGACGGCGAA  TGGCCTATGG

5290        5300        5310        5320        5330        5340
TGTCCGCCTT  TCTCCCTTCG  GGAAGCGTGG  CGCTTTCTCA  ATGCTCACGC  TGTAGGTATC
ACAGGCGGAA  AGAGGGAAGC  CCTTCGCACC  GCGAAAGAGT  TACGAGTGCG  ACATCCATAG

5350        5360        5370        5380        5390        5400
TCAGTTCGGT  GTAGGTCGGT  CGCTCCAAGC  TGGGCTGTGT  CCCGTTCAGC
AGTCAAGCCA  CATTCCAGCAA GCGAGGTTCG  ACCCGACACA  GGGCAAGTCG
```

Figure 19J

(SEQ ID NO.: 23 – Primary Sequence)
(SEQ ID NO.: 29 – Complement)

pD17-hG1b

```
5410                5420                5430                5440                5450                5460
CCGACCGCTG          CGCCTTATCC          GGTAACTATC          GTCTTGAGTC          CAACCCGGTA          AGACACGACT
GGCTGGCGAC          GCGGAATAGG          CCATTGATAG          CAGAACTCAG          GTTGGGCCAT          TCTGTGCTGA

5470                5480                5490                5500                5510                5520
TATCGCCACT          GGCAGCAGCC          ACTGGTAACA          GGATTAGCAG          AGCGAGGTAT          GTAGGCGGTG
ATAGCGGGTGA         CCGTCGTCGG          TGACCATTGT          CCTAATCGTC          TGCGTCCATA          CATCCGGCCAC

5530                5540                5550                5560                5570                5580
CTACAGAGTT          CTTGAAGTGG          TGGCCTAACT          ACGGCTACAC          TAGAAGGACA          GTTATTTGGTA
GATGTCTCAA          GAACTTCACC          ACGGATTGA           TGCCGATGTG          ATCTTCCTGT          CATAAAACCAT

5590                5600                5610                5620                5630                5640
TCTGCGCTCT          GCTGAAGCCA          GTTACCTTCG          GAAAAAGAGT          TGGTAGCTCT          TGATCCGGCA
AGACGCGAGA          CGACTTCGGT          CAATGGAAGC          CTTTTTCTCA          ACCATCGAGA          ACTAGGCCGT

5650                5660                5670                5680                5690                5700
AACAAAACCAC         CGCTGGGTAGC         GGTGGTTTTT          TTGTTTGCAA          GCAGCAGATT          ACGGCGAGAA
TTGTTTTGGTG         GCGACCCATCG         CCACCAAAAA          AACAAACGTT          CGTCGTCTAA          TGCGGCGTCTT

5710                5720                5730                5740                5750                5760
AAAAGGATC           TCAAGAAGAT          CCTTTGATCT          TTTCTACGGG          GTCTGACGCT          CAGTGGAACG
TTTTTCCTAG          AGTTCTTCTA          GGAAACTAGA          AAAGATGCCC          CAGACTGCGA          GTCACCTTGC

5770                5780                5790                5800                5810                5820
AAAACTCACG          TTAAGGGATT          TTGGTCATGA          GATTATCAAA          ATATGAGTAA          ACCTAGATCC
TTTTGAGTGC          AATTCCCTAA          AACCAGTACT          CTAATAGTTT          TATACTCATT          TGGATCTAGG

5830                5840                5850                5860                5870                5880
TTTTAAATTA          AAAATGAAGT          TTTAAATCAA          TCTAAAGTAT          ATTAGAGTAA          ACTTGGTCTG
AAAATTTAAT          TTTTACTTCA          AAATTTAGTT          AGATTTCATA          TAATCTCATT          TGAACCAGAC

5890                5900                5910                5920                5930                5940
ACAGTTACCA          ATGCTTAATC          AGTGAGGCAC          CTATCTCAGC          GATCTGTCTA          TTTCGTTCAT
TGTCAATGGT          TACGAATTAG          TCACTCCGTG          GATAGAGTCG          CTAGACAGAT          AAAGCAAGTA

5950                5960                5970                5980                5990                6000
CCATAGTTGC          CTGACTCCCC          GTCGTGTAGA          TAACTACGAT          ACGGGAGGGC          TTACCATCTG
GGTATCAACG          GACTGAGGGG          CAGCACATCT          ATTGATGCTA          TGCCCTCCCG          AATGGTAGAC
```

Figure 19K
(SEQ ID NO.: 23 – Primary Sequence)
(SEQ ID NO.: 29 – Complement)

pD17-hG1'b

```
6010                6020                6030                6040                6050                6060
GCCCCAGTGC  TGCAATGATA  CCGCGAGACC  CACGCGTCACC  GGCTCCAGAT  TTATCAGCAA
CGGGGTCACG  ACGTTACTAT  GGCGCTCTGG  GTGCGCAGTGG  CCGAGGTCTA  AATAGTCGTT

6070                6080                6090                6100                6110                6120
TAAACCAGCC  AGCCGGAAGG  GCCGAGCGCA  GAAGTGGTCC  TGCAACTTTA  TCCGCCTCCA
ATTTGGTCGG  TCGGCCTTCC  CGGCTCGCGT  CTTCACCAGG  ACGTTGAAAT  AGGCGGAGGT

6130                6140                6150                6160                6170                6180
TCCAGTCTAT  TAATTGTTGC  CGGGGAAGCTA  GAGTAAGTAG  TTTGCGCCAGTT  AATAGTTTGC
AGGTCAGATA  ATTAACAACG  GCCCTTCGAT  CTCATTCATC  AAGCGGTCAA  TTATCAAACG

6190                6200                6210                6220                6230                6240
GCAACGTTGT  TGCCATTGCT  ACAGGCATCG  TGGTGTCACG  CTCGTCGTTT  GGTATGGCTT
CGTTGCAACA  ACGGTAACGA  TGTCCGTAGC  ACCACAGTGC  GAGCAGCAAA  CCATACCGAA

6250                6260                6270                6280                6290                6300
CATTCAGCTC  CGGTTCCCAA  CGATCAAGGC  GAGTTACATG  ATCCGCCATG  TTGTGCAAAA
GTAAGTCGAG  GCCAAGGGTT  GCTAGTTCCG  CTCAAATGTAC  TAGGCGGTAC  AACACGTTTT

6310                6320                6330                6340                6350                6360
AAGCGGGTTAG  CTCCTTCGGT  CCTCCGATCG  TTGTCAGAAG  TAAGTTGGCC  GCAGTGTTAT
TTCGCCCAATC  GAGGAAGCCA  GGAGGCTAGC  AACAGTCTTC  ATTCAACCGG  CGTCACAATA

6370                6380                6390                6400                6410                6420
CACTCANGGT  TATGGCAGCA  CTGCATAATT  CTCTTACTGT  CATGCCATCC  GTAAGATGCT
GTGAGTACCA  ATACCGTCGT  GACGTATTAA  GAGAATGACA  GTACGGTAGG  CATTCTACGA

6430                6440                6450                6460                6470                6480
TTTCTGTGAC  TGGTGAGTAC  TCAACCAAGT  CATTCTGAGA  ATAGTGTTATG  CGGCGACCGA
AAAGACACTG  ACCACTCATG  AGTTGGTTCA  GTAAGACTCT  TATCACATAC  GCCGCTGGCT

6490                6500                6510                6520                6530                6540
GTTGGCTCTTG  CCCGGGCGTCA  ATACGGGATA  ACATACGCGCC  ACTTTTAAAG  ACTTTTAAAG
CAACGAGAAC  GGGCCGCGAGT  TATGCCCTAT  TGTATGCGCGG  TGAAATTTTC  TGAAATTTTC

6550                6560                6570                6580                6590                6600
TGCTCATCAT  TGGAAAAAGT  TCTTCGGGGC  GAAAACTCTC  AAGGATCTTA  CCGGTGTTGA
ACGAGTAGTA  ACCTTTTTGCA  AGAAGCCCCG  CTTTTGAGAG  TTCCTAGAAT  GGCGACAACT
```

Figure 19L
(SEQ ID NO.: 23 – Primary Sequence)
(SEQ ID NO.: 29 – Complement)

pD17-hG1b

```
6610                6620                6630                6640                6650                6660
GATCCAGTTC  GATGTAACCC  ACTCGTGCAC  CCAACTGATC  TTCAGCATCT  TTTACTTTCA
CTAGGTCAAG  CTACATTGGG  TGAGCACGTG  GGTTGACTAG  AAGTCGTAGA  AAATGAAAGT

6670                6680                6690                6700                6710                6720
CCAGCGTTTC  TGGGGTGAGCA  AAAACAGGAA  GGCAAAATGC  CGCAAAAAAG  GGAATAAAGGG
GGTCGCAAAG  ACCCACTCGT  TTTTGTCCTT  CCGTTTTACG  GCGTTTTTTC  CCTTATTTCCC

6730                6740                6750                6760                6770                6780
CGACACGGAA  ATGTTGAATA  CTCATACTCT  TCCTTTTTCA  ATATTATTGA  AGCATTTATC
GCTGTGCCTT  TACAACTTAT  GAGTATGAGA  AGGAAAAAGT  TATAATAACT  TCGTAAATAG

6790                6800                6810                6820                6830                6840
AGGGTTATTG  TCTCATGAGC  GGATACATAT  TTGAATGTAT  TTAGAAAAAT  AAACAAATAG
TCCCAATAAC  AGAGTACTCG  CCTATGTATA  AACTTACATA  AATCTTTTTA  TTTGTTTATC

6850                6860                6870                6880                6890                6900
GGGTTCCGCG  CACATTTCCC  CGAAAAGTGC  CACCTGACGT  CGACGGATCG  GGAGATCTGC
CCCAAGGCGC  GTGTAAAGGG  GCTTTTCACG  GTGGACTGCA  GCTGCCTAGC  CCTCTAGACG

6910                6920                6930                6940                6950                6960
TAGGTGACCT  GAGGCGCGCC  GGCTTCGAAT  AGCCAGAGTA  ACCTTTTTTT  TTTAATTTTAT
ATCCACTGGA  CTCCGCGCGG  CCGAAGCTTA  TCGGTCTCAT  TGGAAAAAAA  AAATTAAAATA

6970                6980                6990                7000                7010                7020
TTTATTTTAT  TTTTGGAGATG  GAGTTTGGCG  CCGATCTCCC  GATCCCCTAT  GGTCGGACTCT
AAATAAAATA  AAAACCTCTAC  CTCAAACCGC  GGCTAGAGGG  CTAGGGGATA  CCAGCCTGAGA

7030                7040                7050                7060                7070                7080
CAGTACAATC  TGCTCTGATG  CCGCATAGTT  AAGCCAGTAT  CTGCTCCCTG  CTTGTGTGTT
GTCATGTTAG  ACGAGACTAC  GGCGTATCAA  TTCGGTCATA  GACGAGGGAC  GAACACACAA

7090                7100                7110                7120                7130                7140
GGAGGTCGGCT  GAGTAGTGCG  CGAGCAAAAT  TTTAAGCTACA  ACAAGGCAAG  GCTTGACCGA
CCTCCAGCGA  CTCATCACGC  GCTCGTTTTA  AAATTCGATGT  TGTTCCGTTC  CGAACTGGCT

7150                7160                7170                7180                7190                7200
CAATTGCATG  AAGAATCTGC  TTAGGGTTAG  GCGTTTTGCG  CTGCTTCGCG  ATGTACGGGC
GTTAACGTAC  TTCTTAGACG  AATCCCAATC  CGCAAAACGC  GACGAAGCGC  TACATGCCCG
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Figure 19M
(SEQ ID NO.: 23 – Primary Sequence)
(SEQ ID NO.: 29 – Complement)

pD17-hG1b

```
           7210       7220       7230       7240       7250       7260
      CAGATATACG CGTTGACATT GATTATTGAC TAGTTATTAA TAGTAATCAA TTACGGGGTC
      GTCTATATGC GCAACTGTAA CTAATAACTG ATCAATAATT ATCATTAGTT AATGCCCCAG

           7270       7280       7290       7300       7310       7320
      ATTAGTTCAT AGCCCATATA TGGAGTTCCG CGTTACATAA CTTACGGGTAA ATGGCCCGCC
      TAATCAAGTA TCGGGTATAT ACCTCAAGGC GCAATGTATT GAATGCCATT TACCGGGGCG

           7330       7340       7350       7360       7370       7380
      TGGCTGACCG CCCAACGACC CCCGCCCATT GACGTCAATA ATGACGTATG TTCCCATAGT
      ACCGACTGGC GGGTTGCTGG GGGCGGGTAA CTGCAGTTAT TACTGCATAC AAGGGTATCA

           7390       7400       7410       7420       7430       7440
      AACGCCAATA GGGACTTTCC ATTGACGTCA ATGGGTGGAC TATTTACGGT AAACTGCCCA
      TTGCGGTTAT CCCTGAAAGG TAACTGCAGT TACCCACCTG ATAAATGCCA TTTGACGGGT

           7450       7460       7470       7480       7490       7500
      CTTGGCAGTA CATCAAGTGT ATCATATGCC AAGTACGCCC CCTATTGACG TCAATGACGG
      GAACCGTCAT GTAGTTCACA TAGTATACGG TTCATGCGGG GGATAACTGC AGTTACTGCC

           7510       7520       7530       7540       7550       7560
      TAAATGGCCC GCCTGGCATT ATGCCCAGTA CATGACCTTA TGGGACTTTC CTACTTGGCA
      ATTTACCGGG CGGACCGTAA TACGGGTCAT GTACTGGAAT ACCCTGAAAG GATGAACCGT

           7570       7580       7590       7600       7610       7620
      GTACATCTAC GTATTAGTCA TCGCTATTAC CATGGTGATG CGGTTTTGGC AGTACATCAA
      CATGTAGATG CATAATCAGT AGCGATAATG GTACCACTAC GCCAAAACCG TCATGTAGTT

           7630       7640       7650       7660       7670       7680
      TGGGCGTGGA TAGCGGTTTG ACTCACGGGG ATTTCCAAGT CTCCACCCCA TTGACGTCAA
      ACCCGCACCT ATCGCCAAAC TGAGTGCCCC TAAAGGTTCA GAGGTGGGGT AACTGCAGTT

           7690       7700       7710       7720       7730       7740
      TGGGAGTTTG TTTTGGCACC AAAATCAACG GGACTTTCCA AAATGTCGTA ACAACTCCGC
      ACCCTCAAAC AAAACCGTGG TTTTAGTTGC CCTGAAAGGT TTTACAGCAT TGTTGAGGCG

           7750       7760       7770       7780       7790       7800
      CCCATTGACG CAAATGGGCG GTAGGCGTGT ACGGTGGGAG GTCTATATAA GCAGAGCTCT
      GGGTAACTGC GTTTACCCGC CATCCGCACA TGCCACCCTC CAGATATATT CGTCTCGAGA
```

Replacement Sheet

Figure 19N
(SEQ ID NO.: 23 – Primary Sequence)
(SEQ ID NO.: 29 – Complement)

pD17-hG1b

```
       7810        7820        7830        7840          7850          7860
CTGGCTAACT AGAGAACCCA CTGCCTTACTG GCTTATCGAA ATTAATACGA CTCACTATAG
GACCGATTGA TCTCTTGGGT GACGAATGAC CGAATAGCTT TAATTATGCT GAGTGATATC

       7870        7880
GGAGACCCAA GCTT
CCTCTGGGTT CGAA
```